

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/921,060A

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DATE 11/02/2001
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#36

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: ANDERSON, DARRELL R.
HANNA, NABIL
LEONARD, JOHN E.
NEWMAN, ROLAND A.
REFF, MITCHELL E.
RASTETTER, WILLIAM H.

(ii) TITLE OF INVENTION: THERAPEUTIC APPLICATION OF CHIMERIC AND
RADIOLABELED ANTIBODIES TO HUMAN B LYMPHOCYTE RESTRICTED
DIFFERENTIATION ANTIGEN FOR THE TREATMENT OF B CELL
LYMPHOMA

(iii) NUMBER OF SEQUENCES: 11

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: PILLSBURY WINTHROP
(B) STREET: 1100 New York Avenue, N.W., Ninth FL.
(C) CITY: Washington
(D) STATE: DC
(E) COUNTRY: USA
(F) ZIP: 20005

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/921,060
(B) FILING DATE: 29-AUG-1997
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Teskin, Robin L.
(B) REGISTRATION NUMBER: 35,030
(C) REFERENCE/DOCKET NUMBER: 037003-0275463

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 202-861-3000
(B) TELEFAX: 202-822-0944

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/921,060ADATE 11/02/2001
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48 (2) INFORMATION FOR SEQ ID NO:1:
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50 (i) SEQUENCE CHARACTERISTICS:
51 (A) LENGTH: 27 base pairs
52 (B) TYPE: nucleic acid
53 (C) STRANDEDNESS: single
54 (D) TOPOLOGY: linear
55
56 (ii) MOLECULE TYPE: DNA (genomic)
57
58
59
60
61 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

62
63 GGGAGCTTGG ATCGATCCTC TATGGTT
64

27

65 (2) INFORMATION FOR SEQ ID NO:2:
66
67 (i) SEQUENCE CHARACTERISTICS:
68 (A) LENGTH: 8540 base pairs
69 (B) TYPE: nucleic acid
70 (C) STRANDEDNESS: single
71 (D) TOPOLOGY: linear
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73 (ii) MOLECULE TYPE: DNA (genomic)
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75
76
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78 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

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83 ATGGTTGCTG ACTAATTGAG ATGCATGCTT TGCATACTTC TGCTGCTGG GGAGCCTGGG 240
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85 GGGGAGCCTG GGGACTTTCC ACACCCTAAC TGACACACAT TCCACAGAAT TAATTCCCCT 360
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87 GTTACATAAC TTACGGTAAA TGGCCCCGCT GGCTGACCGC CCAACGACCC CCGCCCATTG 480
88 ACGTCAATAA TGACGTATGT TCCCATAGTA ACGCCAATAG GGACTTTCCA TTGACGTCAA 540
89 TGGGTGGACT ATTTACGGTA AACTGCCCAC TTGGCAGTAC ATCAAGTGTA TCATATGCCA 600
90 AGTACGCCCC CTATTGACGT CAATGACGGT AAATGGCCCC CCTGGCATTG TGCCCAGTAC 660
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92 ATGGTGATGC GGT TTTGGCA GTACATCAAT GGGCGTGGAT AGCGGTTTGA CTCACGGGGA 780
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94 GACTTTCCAA AATGTCGTAA CAACTCCGCC CAGAGCTGGG TACGTGAACC GTCAGATCGC CTGGAGACGC 900
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218 AACCCACTCG TGCACCCAAC TGATCTTCAG CATCTTTTAC TTTCACCAGC GTTTCTGGGT 8340
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220 GAATACTCAT ACTCTTCCTT TTCAATATT ATTGAAGCAT TTATCAGGGT TATTGTCTCA 8460
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(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 9209 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

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242 ATGGTTGCTG ACTAATTGAG ATGCATGCTT TGCATACTTC TGCCTGCTGG GGAGCCTGGG 240
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Original Text